

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In Re Patent Application of:)
Eric C. Hannah) Examiner: Not Yet Assigned
Application No.: Not Yet Assigned) Art Unit: Not Yet Assigned
Filed: Concurrently Herewith)
For: FAST SECONDARY STRUCTURE)
DISCOVERY METHOD FOR PROTEIN)
FOLDING)

Commissioner for Patents

P.O. Box 1450
Alexandria, Virginia 22313-1450

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Sir:

Enclosed is a copy of Information Disclosure Citation Form PTO-1449 or PTO/SB/08. All references listed on the enclosed Form PTO-1449 or PTO/SB/08 were previously identified in the parent application of the present application, and copies of the references were furnished at that time. It is respectfully requested that the cited documents be considered and that the enclosed copy of Information Disclosure Citation Form PTO-1449 or PTO/SB/08 be initialed by the Examiner to indicate such consideration and a copy thereof returned to applicant(s).

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Express Mail No. EV277415438USDate of Deposit: October 23, 2003Name of Person Mailing Correspondence: Debbie PeloquinDebbie Peloquin
SignatureOctober 23, 2003
Date

Pursuant to 37 C.F.R. § 1.97, the submission of this Information Disclosure Statement is not to be construed as a representation that a search has been made and is not to be construed as an admission that the information cited in this statement is material to patentability.

Pursuant to 37 C.F.R. § 1.97, this Information Disclosure Statement is being submitted under one of the following (as indicated by an "X" to the left of the appropriate paragraph):

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 A statement pursuant to 37 C.F.R. §1.97(e) or

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If there are any additional charges, please charge Deposit Account No. 02-2666.

Respectfully submitted,

BLAKELY, SOKOLOFF, TAYLOR & ZAFMAN LLP

Dated: October 23, 2003


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INFORMATION DISCLOSURE STATEMENT BY APPLICANT

(use as many sheets as necessary)

Complete if Known

Application Number

Filing Date

First Named Inventor: Eric C. Hannah

Art Unit

Examiner Name

Sheet

2

of

2

Attorney Docket Number

42390P11816D

NON PATENT LITERATURE DOCUMENTS

Examiner Initials*	Cite No ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published	T ²
		Al-Lazikani, Bissan et al., "Protein Structure Prediction", Curr. Opin. In Chem. Biol., 5:51-56, 2001	
		Bonneau and Baker, "Ab Initio Protein Structure Prediction: Progress and Prospects", Annu. Rev. Biophys. Struct., 30:73-189, 2001	
		Clementi, Cecilia et al., "Topological and Energetic Factors: What Determines the Structural Details of the Transition State Ensemble and "En-route" Intermediates for Protein Folding? An Investigation for Small Globular Proteins", J. Mol. Biol., 298:937-953, 2000	
		Debe, Derek A. et al., "The Topomer-sampling model of protein folding", Proc. Natl. Acad. Sci USA, 96:2596-2601, March 1999	
		Debe and Goddard III, "First Principles Prediction of Protein Folding Rates", J. Mol. Biol.. 294:619-625, 1999	
		Eyrich, Volker A. et al., "Prediction of Protein Tertiary Structure to Low Resolution: Performance for a Large and Structurally Diverse Test Set", J. Mol. Biol. 288:725-742, 1999	
		Mirny, Leonid A. et al., "Statistical significance of protein structure prediction by threading", PNAS, 97(18) 9978-9983, August 29, 2000	
		Pillardy, Jaroslaw et al., "Recent improvements in prediction of protein structure by global optimization of a potential energy function", PNAS 98(5) 2329-2333, February 27, 2001	
		Thomas and Dill, "An interactive method for extracting energy-like quantities from protein structures", Proc. Natl. Acad. Sci USA 93:11628-11633, October 1996	

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